IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re the Application of)
)
Hugh Gramajo et al.)
)
Serial No. 10/045,612)
)
Filed: October 23, 2001)
)
For: ANTIBIOTIC PRODUCTION II)

SUBMISSION OF SEQUENCE LISTING UNDER 37 C.F.R. §§1.821-1.825 AND PRELIMINARY AMENDMENT

The present submission is in response to the Office communication dated March 14, 2002 enclosing a Notice To File Missing Parts Of Nonprovisional Application.

To comply with the requirements under 37 C.F.R. §§1.821-1.825, submitted herewith is a sequence listing of the amino acids and nucleotides presented in the above-referenced application. The sequence listing is being submitted in both paper copy and computer-readable form. Applicants respectfully request entry of the sequence listing into the above identified patent application. The undersigned hereby verifies that the paper copy and computer readable form of the sequence listing are identical and do not contain any new matter. In the event that a fee is required, the Commissioner is authorized to charge the account of the undersigned, Account No. 04-1406. A duplicate copy of this sheet is enclosed.

Respectfully submitted,

DANN, DORFMAN, HERRELL AND SKILLMAN
A Professional Corporation

By Patrick J. Hagan
Patrick J. Hagan
PTO Registration No. 27,643

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#3

SEQUENCE LISTING

<160> Gramajo, Hugo C
Rodriguez, Eduardo J

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ccctactacg cggccgaacg cggtctggtc gacgacgtca tcgaccccgc cgaaacccgc    1500
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aagcacggca acccgccgca gtga                                           1584
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<212> DNA
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<400> 30

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ggcggcggcc gcgcccgcgc cggctggcgc cgcctggaac gcgagccggg cttccgcgcc     180
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<400> 31

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Arg Val Ala Glu Leu His Gly Ile Arg Ala Ala Ala Leu Ala Gly Pro
            20                  25                  30

Ser Glu Lys Ala Thr Ala Ala Gln His Ala Lys Gly Lys Leu Thr Ala
        35                  40                  45

Arg Glu Arg Ile Glu Leu Leu Leu Asp Pro Gly Ser Phe Arg Glu Val
    50                  55                  60

Glu Gln Leu Arg Arg His Arg Ala Thr Gly Phe Gly Leu Glu Ala Lys
65                  70                  75                  80

Lys Pro Tyr Thr Asp Gly Val Ile Thr Gly Trp Gly Thr Val Glu Gly
                85                  90                  95

Arg Thr Val Phe Val Tyr Ala His Asp Phe Arg Ile Phe Gly Gly Ala
            100                 105                 110

Leu Gly Glu Ala His Ala Thr Lys Ile His Lys Ile Met Asp Met Ala
        115                 120                 125

Ile Ala Ala Gly Ala Pro Leu Val Ser Leu Asn Asp Gly Ala Gly Ala
    130                 135                 140

Arg Ile Gln Glu Gly Val Ser Ala Leu Ala Gly Tyr Gly Gly Ile Phe
145                 150                 155                 160
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Gln Arg Asn Thr Lys Ala Ser Gly Val Ile Pro Gln Ile Ser Val Met
                165                 170                 175

Leu Gly Pro Cys Ala Gly Gly Ala Ala Tyr Ser Pro Ala Leu Thr Asp
            180                 185                 190

Phe Val Phe Met Val Arg Asp Thr Ser Gln Met Phe Ile Thr Gly Pro
        195                 200                 205

Asp Val Val Lys Ala Val Thr Gly Glu Glu Ile Thr Gln Asn Gly Leu
    210                 215                 220

Gly Gly Ala Asp Val His Ala Glu Thr Ser Gly Val Cys His Phe Ala
225                 230                 235                 240

Tyr Asp Asp Glu Glu Thr Cys Leu Ala Glu Val Arg Tyr Leu Leu Ser
                245                 250                 255

Leu Leu Pro Gln Asn Asn Arg Glu Asn Pro Pro Arg Ala Glu Ser Ser
            260                 265                 270

Asp Pro Val Asp Arg Arg Ser Asp Thr Leu Leu Asp Leu Val Pro Ala
        275                 280                 285

Asp Gly Asn Arg Pro Tyr Asp Met Thr Lys Val Ile Glu Glu Leu Val
    290                 295                 300

Asp Glu Gly Glu Tyr Leu Glu Val His Glu Arg Trp Ala Arg Asn Ile
305                 310                 315                 320

Ile Cys Ala Leu Ala Arg Leu Asp Gly Arg Val Val Gly Ile Val Ala
                325                 330                 335

Asn Gln Pro Gln Ala Leu Ala Gly Val Leu Asp Ile Glu Ala Ser Glu
            340                 345                 350

Lys Ala Ala Arg Phe Val Gln Met Cys Asp Ala Phe Asn Ile Pro Ile
        355                 360                 365

Ile Thr Leu Leu Asp Val Pro Gly Phe Leu Pro Gly Val Asp Gln Glu
    370                 375                 380

His Gly Gly Ile Ile Arg His Gly Ala Lys Leu Leu Tyr Ala Tyr Cys
385                 390                 395                 400

Asn Ala Thr Val Pro Arg Ile Ser Leu Ile Leu Arg Lys Ala Tyr Gly
                405                 410                 415

Gly Ala Tyr Ile Val Met Asp Ser Gln Ser Ile Gly Ala Asp Leu Thr
            420                 425                 430

Tyr Ala Trp Pro Thr Asn Glu Ile Ala Val Met Gly Ala Glu Gly Ala
        435                 440                 445

Ala Asn Val Ile Phe Arg Arg Gln Ile Ala Asp Ala Glu Asp Pro Glu
    450                 455                 460

Ala Met Arg Ala Arg Met Val Lys Glu Tyr Lys Ser Glu Leu Met His
465                 470                 475                 480

Pro Tyr Tyr Ala Ala Glu Arg Gly Leu Val Asp Asp Val Ile Asp Pro
                485                 490                 495

Ala Glu Thr Arg Glu Val Leu Ile Thr Ser Leu Ala Met Leu His Thr
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Lys His Ala Asp Leu Pro Ser Arg Lys His Gly Asn Pro Pro Gln
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<210> 32
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<212> PRT
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<400> 32
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Glu Val Ala Ala Ile Thr Ala Leu Leu Leu Ala Arg Ala Ala Ala Arg
20 25 30

Pro Ala Glu Ile Ala Pro Thr His Gly Gly Gly Arg Ala Arg Ala Gly
35 40 45

Trp Arg Arg Leu Glu Arg Glu Pro Gly Phe Arg Ala Pro His Ser Trp
50 55 60

Arg
65